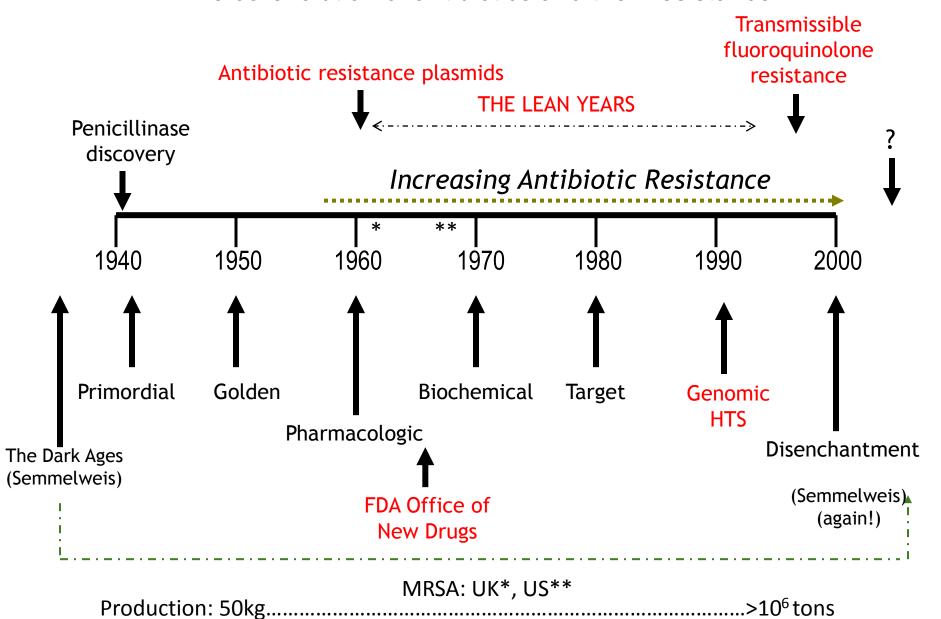
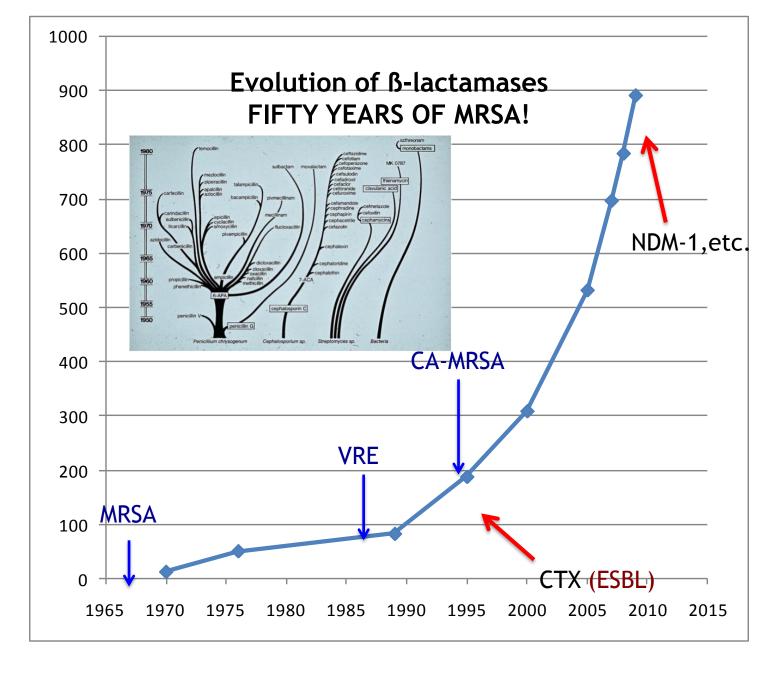


ATA, Paris, 25/09/2012

#### The co-evolution of antibiotics and their resistance





# The most expensive hydrolytic reaction in history!

### Fifty years of "Industrial Research" on Antibiotics

#### **Environmental**

Doubling time: months

Aeration: low

Nutrients: variable

Carbohydrate:

limited

Water: variable

Temperature: 0-40

pH: 2-10

Yield: µgrams

#### **Production**

Doubling time:

hours

Aeration: high

\*Nutrients: high and

constant

Carbohydrate: high

Water: unlimited

Temperature: 25-30

pH: 6-7

Yield: grams

#### \*Typical substrates for fermentation

Molasses (blackstrap)

Fish Meal (herring, anchovy)

Citrus pulp

Asparagus juice

Cottonseed oil

Malt extract

Beef extract

Bovine blood

Pork liver

Distillers solubles

Hydrolysed rabbit fur

#### We Live In A Microbial World

Microbial viruses on the Earth: 1x10<sup>31</sup>

Microbes on the Earth: 5x10<sup>30</sup>

Stars in the Universe: 7x10<sup>21</sup>

Humans on the Earth: 6x109

Human genes in one person: 2.5x10<sup>4</sup>

Human cells in one person: 1x10<sup>13</sup>

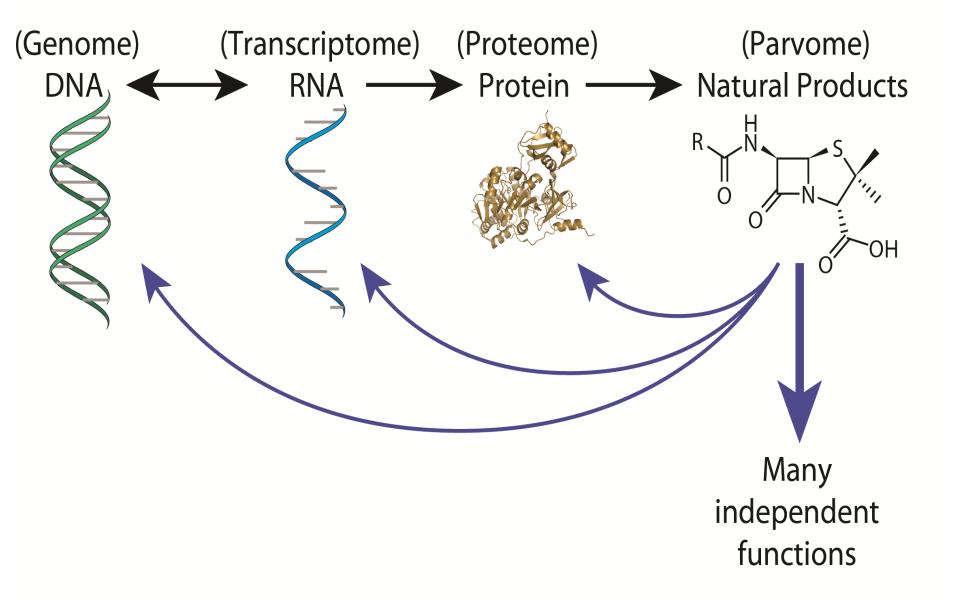
Microbial genes in human gut: 3x10<sup>6</sup>

Microbial cells in human gut: 1x10<sup>14</sup>

Bubbles in one bottle of champagne: 1x10<sup>5</sup>

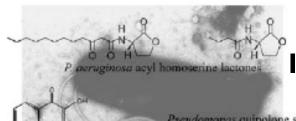
We are less than 10% of what we think we are!

## The "Revised" Central Dogma



"Life would not exist with macromolecules alone." Stuart Schreiber, 2005

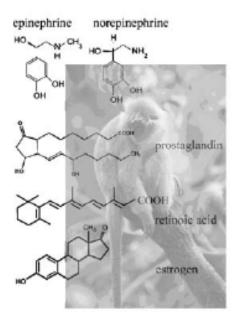
#### MONERA

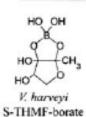


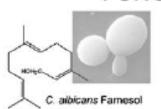
#### **Inter-Kingdom Signalling**

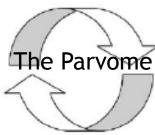
Pseudomonas quinolone signal

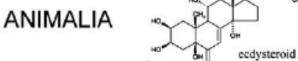
#### **FUNGI**



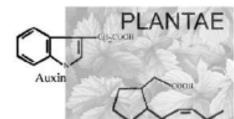








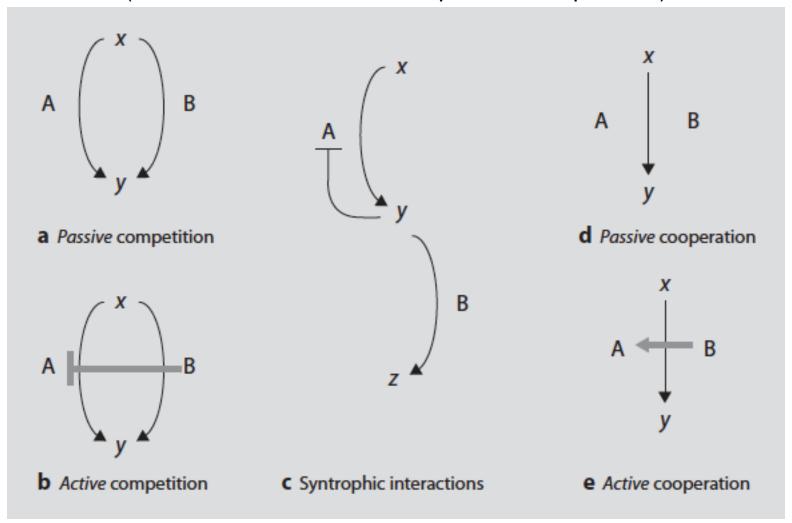
C. elegans 3-Keto-4-cholestenoic acid



(Rumbaugh, 2007)

Jasmonic Acid

# Signaling Interactions between Cells in Nature (Communication, Cues, Competition, Cooperation)



# Prospects for natural product drug discovery

- Bioactivity-based screens
- Combinatorial synthesis and biosynthesis
- Chemical library screening by "docking"
- Co-culturing producing strains: pathway activation
- Genomic identification of biosynthetic pathways
- Heterologous expression of pathways
- Drug combinations
- New approaches to vaccines
- A "new" look at plants and other sources
- Personalized treatments

# Genomics and Drug Discovery

(A new paradigm)

Genome or Metagenome sequence



Bioinformatic scanning for biosynthetic gene clusters



Prediction of structure or molecule class



Virtual docking for target identification



Compound production by cluster expression in "designer" host

# Analysis of molecular interactions in bacterial cell networks, using high-resolution MS with nanoDesi probe (Watrous *et al* 2012)

